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In the specification:

Please replace Table III on pages 89-91 with Table III as follows:

TABLE III

SEQ ID NO. 1

>40_TCC_13F11_M13F.fa TIME: Sun Sep 10 11:42:06 2000 trimming information: raw_sequence:582 (high quality:29-320) sequence:97-252

[length:156]
TCCGTCTCATTGAGGGTCCTGAGGAAGTTGATCTCATCATTCAGGGCATC
CACCTTGGCCTCCAGCTCCACCTTGCTCATGTAGGCAGCATCCACATCCT
TCTTCAGCACCACAAACTCATTCTCAGCAGCTGTGCGGCGGTTAATTTCA

TCTTCG

SEQ ID NO. 2

SEO ID NO. 3

>20_TCC_60H4_M13F.TXT.fa ,constant: -1, poly A: no CANTATATAACNAATTGGAGCTCAATNGCNCGCGGNCGCGTGTCTTCTGGGTAGAGGAT GNGAAGGAAGGACCCTTACCCCCGGCTCTTCTCCTGACCTGCCAATAAAAATTTATGGT CCAAGGNAAAANA

SEQ ID NO. 4

>26_TCC_44C1_M13F.TXT.fa ,constant: -1, poly A: no
ACTCATTGAACTTGAGCTCCGANTCCTGATTCNCATCNAAGCTCTNNATCTGCTCATCAN
GAGANCCCACATCCTTGAGCAGATGGNGCANCTGCTGNTTAACCANCTCTNNGAACTCGN
AGANNNTAAGGCTATCCTTCCGGNCCTCCTGCCTTGCAAAGGTGAAGAAAGTGGTGNNCA
CNGTCNCAATGGANTCCTCTAGCTCTGTCAGTGGTTCTGCTGCNATTATGGAACCTGAGG
CCAAAGCTGATGTCCTCAAGGGGCTAGCTGACCTTTGTCAGGGCTGACCTCTCCTCAGCG
GCAGCAGGGCAGAGTGCTGAACCCAGGAACCCACAGATCCTCCCCGNTCCTGTCTCCCGG
TGACAAGGGTCCTGGAACGGGGCGTCTCTGACTCCCTGCTCCAGGGCTTTAAGT

SEQ ID NO. 5
>29_TCC_48G1_M13F.TXT.fa ,constant: -1, poly A: no
ACTTTGAGAAGGCAGGACTCAAATGATGCCCTGGAGATGTCACAGATTCCTGGCAGAGCC
ATGGTCCCAGGCTTCCCAAAAGTGTTTGTTGGCAATTATTCCCCTAGGCTGAGCCTGCTC
ATGT

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SEQ ID NO. 6

>31_TCC_65B9_M13F.TXT.fa ,constant: -1, poly A: yes GACTAGAACCCACCCCTTNCCTTCCAGCCTTTCTGTCATCATCTCCACAGNCCANCCAT CCCCTGAGCACACCATCTCATGCAGGCCCCACCTGCCAATAGTAATAAAGCAATGT CACTTTGTTAAAACATGAAAAAAAAA

SEQ ID NO. 7

SEQ ID NO. 8

>10 TCC 53H11 T3

SEQ ID NO. 9

>07_TCC_57B3_M13F.TXT ,constant: -1, poly A: yes GGTACCGACGGACCTGCGGAGACTCCTGCCCTGTTGTGTATAGATGCAAGATATTTATAT ATATTTTTGGTTGCAATATTAAATACAGACACTAAGTTATAGTATATCTGGCAAGCCAAC TTGTAAATCACCACCTCACTCCTGTACTTACCTAAACAGATATAAATGGCTGGTTTTTAA GAAAAAAAA

SEQ ID NO. 10

>11_TCC_25F2_M13F.TXT ,constant: -1, poly A: no ACCCTGGGAGAGAGTTTGAAGAAACCACAGCTGATGGCAGAAAACTCAGACTGCTGCA ACTTTACAGATGGTGCATTGNGTCAGCATAGGAGTGAGATGGGGAAAGCACANTAA CAAGAAAATTGANAGATGNTAAATTAGTGNTGGAGTGTCATGAACAATGCACCTGT

SEO ID NO. 11

>25_TCC_50G5_M13F.TXT ,constant: 17, poly A: yes
TAGTGTGGAAGCATAGTGAACACACTGATTAGGTTATGGTTTAATGTTACAACAACTATT
TTTTAAGAAAAACATGTTTTAGAAATTTGGTTTCAAGTGACATGTGTGAAAACAATATCG
ATACTACCATAGTGAGCCATGATTTTCTAAAAAAAAA

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SEQ ID NO. 12

>26_TCC_50G6_M13F.TXT ,constant: 17, poly A: yes
TAGTGTGGAAGCATAGTGAACACACTGATTAGGTTATGGTTTAATGTTACAACAACTATT
TTTTAAGAAAAACAAGTTTTAGAAATTTGGTTCAAGTGACATGTGTGAAAAACAATATTGT
ATACTACCATAGTGAGCCATGATTTTCTAAAAAAAAA

SEQ ID NO. 13

SEQ ID NO. 14

SEQ ID NO. 15

>38_TCC_56E11_M13F.TXT ,constant: -1, poly A: yes CTCTCCAGTTTGCACCTGTCCCCACCCTCCACTCAGCTGTCCTGCAGCAAACACTCCACC CTCCACCTTCCATTTTCCCCCACTACTGCAGCACCTCCAGGCCTGTTGCTATAGAGCCTA CCTGATGTCAATAAACAACAGCTGAAGCAAAAAAAAA

SEQ ID NO. 16

SEQ ID NO. 17

>53_TCC_79G2_M13F_E07_054.abl.TXT ,constant: 16, poly A: no TGTCCGTCTTCACCCATCCCCAAGCCTACTAGAGCAAGAAACCAGTTGTAATATAAAATG CACTGCCCTACTGTTGTATGACTACCGTTACCTACTGTTGTCATTGTTATTACAGCTAT GGCCACTATTATTAAAGAGCTGTGTAACATCAAAAAAA

SEQ ID NO. 18

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SEQ ID NO. 19

>35_TCC_21D6_M13F_C05_037.abl.fa TIME: Wed Aug 9 12:48:31 2000 trimming information: raw_sequence:889 (high quality:34-340) sequence:95-456 [length:362]

CTTTGACGTGGAGAGGAACTCCTGCAATAACTTCATCTATGGAGGCTGCC
GGGGCAATAAGAACAGCTACCGCTCTGAGGAGGCCTGCATGCTCCGCTGC
TTCCGCCAGCAGGAGAATCCTCCCCTGCCCCTTGGCTCAAAGGTGGTGCT
TCTGGCGGGGGCTGTTCGTGATGGTGTTGATCCTCTTCCTGGGAGCCTCCA
TGGTCTACCTGATCCGGGTGGCACGGAGGAACCAGGAGCGTGCCCTGCGC
ACCGTCTGGAGCTCCGNAGATGACAAGGAGCAGCTGGTGAAGAACACATA
TGTCCTGTGACCGCCCTGTCGCCAAGAGGACTGGNGAAAGGGAGGGAGA
CTATGTGTGAGC

BL

SEQ ID NO. 20

>46_TCC_27H5_M13F_F06_058.ab1.fa TIME: Wed Aug 9 12:48:35 2000 trimming information: raw_sequence:892 (high quality:169-406) sequence:170-287 [length:118]

AAAAAGAGTAAAACACTTTCAGTTTCTCCCCTTTAGCCCCTAAAACAACA TCTTACAGTCTGGATCTGGATCTACCTATACAGTCCTACATTAGCTTCTA

AAATATTTGTCAGGAGGG-

Please replace Table IV on pages 92-94 with Table IV as follows:

TABLE IV

SEQ ID NO. 21

>31_TCC_10E8_M13F.fa TIME: Sun Sep 10 11:42:01 2000 trimming information: raw_sequence:549 (high quality:25-313) sequence:98-313 [length:216]

CCCAAATGGAATGTTGCCCCCTTAAACACCATTTTCCCTCCAGGACCACC TTGGTTTCTAGGCACTGTGGTTCTTGGCAGGGGCTGTCTTAGGTAAAAGG GTAGTTGTGGAGCTACAGTCTGAAGAACATAGCTTGGGCTCAAGTTCAAA TGAGCCATCTTTTTCCTTTGCGTTTTTCTTGACTGAAGGTGAGATGTTAT TTGTGGCATGTGAACT

SEQ ID NO. 22

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SEQ ID NO. 23

>11 TCC_101E11_M13F.TXT.fa ,constant: 15, poly A: yes

GCAATAAAGCTGTCCATTCAATTCCAAATACTGGTTTTAAGNGTATAGCCACTGATATTC TTTCATGTNTAGAAATTCTTTCTGTTATTATTCAAGAAAATGTTTTTAATCATGCTAATA AACTTTTTTGGAGATGAAAAAAAA

SEQ ID NO. 24

>15_TCC_57C3_M13F.TXT.fa ,constant: -1, poly A: no

GGNACCACGTACCTGCTGAATGTNTCNNCGNNATGNCGNCAGGCCATGCTGTTGCTGATN TANTACTNTGAAAATANGGATATCATGATGGGAATGCATGTCATGAGGTCCAGANTCGTT CTACTGTCNATAANCTGTNTACTNGCGTTGANAANAAANGATGTCAAAGNCCCCCCGTAA AAANGTA

SEQ ID NO. 25

>44_TCC_70E8_M13F.TXT.fa ,constant: 15, poly A: yes

SEQ ID NO. 26

>03 TCC_57E11_T7.TXT.fa ,constant: 16, poly A: yes

SEQ ID NO. 27

>08 TCC_70E7_M13F_H01_015.ab1.TXT ,constant: -1, poly A: no

GGATCGACGACCTGCTTCCCAGANGCGNNCNNGAGGNCCNCTTGTTNNNGNCNNGNANAC NNACCCANTTNANTTNNAGCCTTTNTGNAATAAATATACACAGGCCACCCATGCCNTGAG CACACTAACCACNTGATGCAGGCCCCACCTTGCCAATAGTAATAAAGCANTGGGACGTTT TTTA

SEQ ID NO. 28

>13 TCC_71E4_M13F_E02_018.ab1.TXT ,constant: -1, poly A: no

GGGCCAAAGCCCGNGCATCCAANCCCANGCAAGGNACAAANGANCNNGGAGAGGANNACCCAAGCANNTNNCAACCATCAAATGGAGGGCANGCCCGGGG

SEO ID NO. 29

>15 TCC_71H8_M13F_G02_019.ab1.TXT ,constant: -1, poly A: no

GGGCCAAAGCCGNGCATCCAANCCCANCGCANGGNANAAANGANGANGGANANGGATNAC CCANGCCTNTATTAACCATCAANTGGGANGGCAAGCCCGGGGCATNTATTGATT

SEQ ID NO. 30

>21 TCC_43E2_M13F.TXT ,constant: -1, poly A: no

AGGACCCCTGAANACNACACAGATCTGTGNGAAACAANGGNACNTAGCGTCCCNAAAGTG CCNGGTTNNNGTANNCNNAGNGNGACCNGNGCNCATNT

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SEQ ID NO. 31

>24_TCC_96C7_M13F.TXT ,constant: 16, poly A: yes

SEQ ID NO. 32

>57_TCC_80C9_M13F_A08_056.ab1.TXT ,constant: 17, poly A: yes CTGCAGGAGTCAGCGTTCAATCTTGACCTTGAAGATGGGAAGGATGTTCTTTTTACGTAC CAATTCTTTTGTCTTTTGATATTAAAAAGAAGTACATGTTCATTGTAGAGAATTTGGAAA CTGTAGAAGAAGAATCAAGAAGAAAAATAAAAATCAGCTGTTGTAATCACCTAGCAAAAAA

SEQ ID NO. 33

>14_TCC_9B6_M13F_F02_026.ab1.fa TIME: Wed Aug 9 12:48:25 2000 trimming information: raw_sequence:871 (high quality:73-413) sequence:98-394 [length:297] CACGCATATGGGGCCAGTTCCACATATTTGGCAACCAGACCAGCATCCAG GACAACACAAAGTATGTTGTTTGTTAGAGGGCTTGGGACATTTCACT CTTTGCCAGCCTCAGCTTAATCCAGGAGACAAAGATTATTTTCCTTATTA TCTCTTCTGCATAGGATCTGCAATCAGAACTATTGAACTTCTCCATTCAG ACCGCCACTCACACCTATGGGAAAAGGGTAATGTATCATCGGCTTAGCAA CAGGGAATACTATTCGTATGATGGAAAATGGGGACAAAAGGCTTTGG

SEQ ID NO. 34

SEQ ID NO. 35

>26_TCC_13H10_M13F_B04_032.ab1.fa TIME: Wed Aug 9 12:48:29 2000 trimming information: raw_sequence:874 (high quality:67-356) sequence:99-261 [length:163] CTAACCCACGATTCTGAGCCCTGAGTATGCCTGGACATTGATGCTAACAT GACCATGCTTGGGATGTCTCTAGCTGGTCTGGGGATAGCTGAGCACTTA CTCAGGTGGCTGAAATGACACCTACGAAGGAATGAGTGCTATAGAGA GGAGAGAGGAGTG



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SEQ ID NO. 36 >28_TCC_16D12_M13F_D04_041.ab1.fa TIME: Wed Aug 9 12:48:29 2000 trimming information: raw sequence:866 (high quality:71-411) sequence:95-602 [length:508] CAGCTGATGTCATGTGGTGCTGAGAAGAAAGCAGATCACACTTCATCACA GAAAGAATGCCTTGTGATTATCTTCTCCACATCTGAAATTCCTTTTGACA CCTGCATTGGGCCGACTGCCATTCCCATGACTGCTGCACCTGCGTTTTTA GAGAATGCCTCATAACCCACTGATTCTCATTCACAGAGAATGGGAATACG GAATGAAGAAAGATTCCAGCAGCTTATAGAAGGATAGCAATATTTTGGGA CAGGGAAAATCCTGTCATACCTCACCTCTTCCTCAGGAGGAGTTCTGAGC TGGTCCTGCTTTTCATAGNTGTTTCTTTTCTTCCACTTAAGAACTCATAG ATTTTTCTTACTGTCCTAAGGAAGTCCTTACCTCTGAGGTATCTCCTCAA TGAATACTGTTTTCAAGGCTGAAATAGTTCATTATGTTAATAACCTTCTT TATGTTCTCAGGGAAATGCTTAGGTGGTGTCACAAAAAGGGCCTTTTCTT **TNCTTTNC**

SEQ ID NO. 37 >29_TCC_17A5_M13F_E04_034.ab1.fa TIME: Wed Aug 9 12:48:30 2000 trimming information: raw_sequence:861 (high quality:83-477) sequence:99-187 [length:89] CTTCAAAAAGTGTATTGTCAAACATACCTAACTTTCTTGCAATAAATGCA AAAGAAACTGGAACTTGACAATTATAAATAGTAATAGTG

SEQ ID NO. 38

>54_TCC_30E5_M13F_F07_062.ab1.fa TIME: Wed Aug 9 12:48:37 2000 trimming information: raw sequence:836 (high quality:65-394) sequence:90-235 [length:146] CAATTTGTTATAGTATAGTATCAAATTTCTATATAGATTTTATACCTCAG TGGGGAAAAATAACTGATTCCAATGACATTCATTTTGTTTTCATCTGTGA TAGTCATGGATGCTTTTATTTTCCTTGGGGTGCTGAAATTGAGCTG

SEO ID NO. 39 >59_TCC_34D5_M13F_C08_065.ab1.fa TIME: Wed Aug 9 12:48:39 2000 trimming information:

raw sequence:875 (high quality:63-434) sequence:96-244 [length:149] CCTGCCAAAATCCTACCACAGGATAACATTACAAGCAAAAAATTTACATG TTCCAAAGTCTACCACACTCAAGAAGTTACTAAGAACTCTTGCAGAATAA AAGTCACCATTTTAGAAATGCAAACCCACTTCCAACCTTTGCACAGTCC

SEO ID NO. 40

>72_TCC_37E11_M13F_H09_079.ab1.fa TIME: Wed Aug 9 12:48:43 2000 trimming information: raw sequence:899 (high quality:35-432) sequence:97-444 [length:348] CATTTTTAGTGACATTTTAAAAGCAGTCAGATTCTATAAATGGCAAGTAA GCCTGAAGTGAGGATACTGCAATTTTCGGAGAAAAGAACAGCAGCTCTTT ACAATTCTTGAACTGATGTCAGCACCCGAGTGGCTCCTGAATTTAAGTCT GGGACGACATCTTTTATTTTTACATGAATCTTTAAACAATTCTGTGAGCA AAGTTTGTAGCTGCTGGATTATTGTCTGTCTTTATAGCAAGTTCCAGTAA ACCACAAGTATGGCAAAGCTTATCCAATTTTATGCTTGNAGCAGTCAG--



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Please replace Table 6 on pages 1-100 with Table 6 as follows:

-Table 6 Polynucleotides corresponding to the Genes described in Table 5.

SEQ ID NO. 41

>17 TCC 70E7 1 M13F.fa

GGTAGACGTACCTGCGTCCCAGACTTGACCAGGTGGATCTCCTGTTTTAC TCACGAGGACTTTCCCAGGAAAACCATGCCACTAGCAAAATAATATAAAC AAAGGA

SEQ ID NO. 42

>17_TCC_70E7_1_M13R.fa

TTTTTTTTTTTTTTTGGCTAGAGGCATGGATATCCTGGGAAAGCTCTCC TGAGTAAAAGACGAGAGACACCTGGTGAAGACTGGAACGCATGTACGTCT ACC

SEO ID NO. 43

>contig_TCC_101E11_RF.fa

SEQ ID NO. 44

>contig_TCC_102C5_RF.fa

SEO ID NO. 45

>contig_TCC_21G7_RF.fa



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SEQ ID NO. 46

>contig_TCC_25F2_RF.fa

CCCTGGGAGAAGTTTGAAGAAACCACAGCTGATGGCAGAAAAACTCAG ACTGTCTGCAACTTTACAGATGGTGCATTGGTTCAGCATCAGGAGTGGGA TGGGAAGGAAAGCACAATAACAAGAAAATTGAAAGATGGGAAATTAGTGG TGGAGTGTGTCATGAACAATGTCACCTG

SEQ ID NO. 47

>contig_TCC_34A5_RF.fa

SEQ ID NO. 48

>contig_TCC_36B5_RF.fa

SEQ ID NO. 49

>contig TCC 42G5_RF.fa

CCTTCCGAAATACTTCCTCCAGGTGGCAGCACCAAGAATATTTCTGGAAG
CATGTGATGAGGTTGTGTGATGAAGATAGAGCCCATTGTGCTGTCTCTCCA
GGACACGTTGTGTGGCGTTGAAGAGCAGAAAGCAATGAAGTCCTTCTCCA
CGTGGGTCTTGTAAACAGCATCTTCCTCCAGGTTCTCAGATGACTGTGAA
GAGGCCACTTCCAAGGATGCTGGAGAGTCTCTGACCCACAGTTCCCACG
GTTTGCACCTCTGCAGGCCTGGACAATGATGACCTTGGGTTTGTCCTTCA
GACTGAGGCAGTTGCGGTTGTTGAATATCTGGAAGATGGTGTCATAAAGC
AGCACATCTGGTTTTTTCTCATCATGCACAGTTCCGCAGATTCCCTCCAG
GATGCCATGAGACATGGG

SEQ ID NO. 50

>contig_TCC_44C1_RF.fa

CTCATTGAACTTGAGCTCCGAGTCCTGATTCACATCCAAGCTCTTCATCT
TCTCATCAAGAGAGCCCACATCCTTGAGCAGATGGGGCAACTGCTGGGTA
ACCAGCTCTTTGAACTCGTTGACGCTGAGGCTATCCTTCCGGCCCTCCTG
CCTTGCAAAGGTGAAGAAGGTGGTGACCACGGTCTCAATGGACTCCTCA
GCTCTGTCAGTGGTTCTGCTGCCATTAGGACCCTGAGGCCAAAGCTGATG
TCCTCAAGGGGCTAGCTGACCTTTGTCAGGGCTGACCTCTCCTCAGCGGC
AGCAGGGCAGAGTGCTGAACCCAGGACCCCACAGATCCTCCCCGGTCCTC
TCTCCCGGTGACAAGGGTCCTGGAACGGGGCGTCTCTGACTCCCCAGCTCCC
AGGACGGGTTTAG



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SEQ ID NO. 51

>contig_TCC_54C11_RF.fa

SEQ ID NO. 52

>contig_TCC_56E11_RF.fa

SEQ ID NO. 53

>contig_TCC_57B3_RF.fa

SEQ ID NO. 54

>contig TCC 58A3_RF.fa

SEQ ID NO. 55

>contig_TCC_70E8_RF.fa

TTTTTTTTTTTTTTTTGCTAGTGGCATGGTTTTCCTGGGAAAGTCCTCC TGAGTAAAAGAGGAGACACCTGGTGAAGACTGGGACGCAGGTACGTCT ACC

SEQ ID NO. 56

>contig_TCC_71E4_RF.fa



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SEQ ID NO. 57

>contig_TCC_75E3_RF.fa

AAAGAGGCGCAGGGGCCTGGAGATCCTCCTGCAGACCACGCCCGTCCT GCCTGTGGCGCCGTCTCCAGGGGCTGCTTCCTCCTGGAAATTGACGAGG GTGTCTTGGGCAGAGCTGGCTCTGAGCGCCTCCATCCAAGGCCAGGTTCT CCGTTAGCTCCTGTGGCCCCACCCTGGGCCTGGGCTGGAATCAGGAATA TTTTCCAAAGAGTGATAGTCTTTTGCTTTTGCAAAACTCTACTTAATCC AATGGGTTTTTCTCTGTACAGTAGATTTTCCAAATGTAATAAACTTTAAT

SEQ ID NO. 58

>contig_TCC_78B11_RF.fa

SEQ ID NO. 59

>contig TCC 89G3_RF.fa

SEQ ID NO 60

>contig TCC_92D7_RF.fa

TTTTTTTTTTTTTGAAGACAACTTTTAGAAACTGATGTTTATTTTCCA TCAACCATTTTTCCATGCTGCTTAAGAGCCTATGCAAGAACAGCTTAAGA CCAGTCAGTGGTTGAAGTC

SEQ ID NO. 61

>contig_TCC_93G5_RF.fa

GACTACCAGACCAACAAAGCCAAGCATGATGAGCTGACCTATTTCTGATC
CTGACTTTGGACAAGGCCCTTCAGCCAGAAGACTGACAAAGTCATCCTCC
GTCTACCAGAGCGTGCACTTGTGATCCTAAAATAAGCTTCATCTCCGGGC
TGTGCCCCTTGGGGTGGAAGGGGCAGGATTCTGCAGCTGCTTTTGCATTT



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The first two sequences are from opposite ends of the same polynucleotide, (and are thus in the same gene). All the other 21 sequences are contigs.

Marked up versions of new Tables III, IV, and 6 showing the changes made are attached hereto as ${\tt Exhibits}$ B, C and D, respectively.

Applicants maintain that the amendments to Tables III, IV, and 6 were made to introduce the sequence ID numbers and raise no issue of new matter. Accordingly, applicants respectfully request that the Amendment be entered.

H

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Sequence Listing

The Notice to Comply With Requirements for Patent Applications Containing Nucleotide Sequence And/Or Amino Acid Sequence Disclosures indicates that a copy of the "Sequence Listing" in computer readable form has not been submitted as required by 37 C.F.R. 1.821(e).

In response, applicants submit as **Exhibit E** hereto a computer readable form (CRF) copy of the "Sequence Listing" as required by 37 C.F.R. §1.825(e) and as **Exhibit F** hereto a courtesy paper copy of the sequence listing. Further, applicants submit herewith a Statement in accordance with 37 C.F.R. §1.821(f) as **Exhibit G**, certifying that the computer readable form containing the nucleic acid and/or amino acid sequences as required by 37 C.F.R. §1.821(e) contains the same information which was submitted as the "Sequence Listing" and contains no new matter.

If a telephone interview would be of assistance in advancing prosecution of the subject application, applicants' undersigned attorney invite the Examiner to telephone him at the number provided below.

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No fee is deemed necessary in connection with the filing of this Communication. If any other fee is required, authorization is hereby given to charge the amount of any such fee to Deposit Account No. 03-3125.

Respectfully submitted,

I hereby certify that this correspondence is being deposited this date with the U.S. Postal Service with sufficient postage as first class mail in an envelope addressed to: Assistant Commissioner For Patents, P.O. Box 2327, Arlington, VA 22202, Box:

phin P. White

eg. No. 28,678

2(1(0

Date

John k. White

Registration No. 28,678
Attorney for Applicants
Cooper & Dunham LLP
1185 Avenue of the Americas
New York, New York 10036
(212) 278-0400

Marked-up Version of the Table III

Additions to the text are indicated by underlining.

TABLE III

SEO ID NO. 1

>40_TCC_13F11_M13F.fa TIME: Sun Sep 10 11:42:06 2000 trimming information: raw_sequence:582 (high quality:29-320) sequence:97-252 [length:156]

TCCGTCTCATTGAGGGTCCTGAGGAAGTTGATCTCATCATCAGGGCATC
CACCTTGGCCTCCAGCTCCACCTTGCTCATGTAGGCAGCATCCACATCCT
TCTTCAGCACCACAAACTCATTCTCAGCAGCTGTGCGGCGGTTAATTTCA
TCTTCG

SEO ID NO. 2

SEO ID NO. 3

>20_TCC_60H4_M13F.TXT.fa ,constant: -1, poly A: no CANTATATAACNAATTGGAGCTCAATNGCNCGCGGNCGCGTGTCTTCTGGGTAGAGGGAT GNGAAGGAAGGAACCCTTACCCCCGGCTCTTCTCCTGACCTGCCAATAAAAATTTATGGT CCAAGGNAAAANA

SEO ID NO. 4

>26_TCC_44C1_M13F.TXT.fa ,constant: -1, poly A: no
ACTCATTGAACTTGAGCTCCGANTCCTGATTCNCATCNAAGCTCTNNATCTGCTCATCAN
GAGANCCCACATCCTTGAGCAGATGGNGCANCTGCTGNTTAACCANCTCTNNGAACTCGN
AGANNNTAAGGCTATCCTTCCGGNCCTCCTGCCTTGCAAAGGTGAAGAAAGTGGTGNNCA
CNGTCNCAATGGANTCCTCTAGCTCTGTCAGTGGTTCTGCTGCNATTATGGAACCTGAGG
CCAAAGCTGATGTCCTCAAGGGGCTAGCTGACCTTTGTCAGGGCTGACCTCTCCTCAGCG
GCAGCAGGGCAGAGTGCTGAACCCAGGAACCCACAGATCCTCCCCGNTCCTGTCTCCCGG
TGACAAGGGTCCTGGAACGGGGCGTCTCTGACTCCCTGCTCCAGGACGGGTTTAAGT

SEO ID NO. 5

>29_TCC_48G1_M13F.TXT.fa ,constant: -1, poly A: no ACTTTGAGAAGGCAGGACTCAAATGATGCCCTGGAGATGTCACAGATTCCTGGCAGAGCC ATGGTCCCAGGCTTCCCAAAAGTGTTTGTTGGCAATTATTCCCCTAGGCTGAGCCTGCTC ATGT

SEO ID NO. 6

>31_TCC_65B9_M13F.TXT.fa ,constant: -1, poly A: yes GACTAGAACCCACCCCTTNCCTTCCAGCCTTTCTGTCATCATCTCCACAGNCCANCCAT CCCCTGAGCACACTAACCATCTCATGCAGGCCCCACCTGCCAATAGTAATAAAGCAATGT CACTTTGTTAAAACATGAAAAAAAA

SEO ID NO. 7

SEO ID NO. 8

>10 TCC 53H11 T3

SEO ID NO. 9

>07_TCC_57B3_M13F.TXT ,constant: -1, poly A: yes GGTACCGACGGACCTGCGGAGACTCCTGCCCTGTTGTGTATAGATGCAAGATATTTATAT ATATTTTTGGTTGCAATATTAAATACAGACACTAAGTTATAGTATATCTGGCAAGCCAAC TTGTAAATCACCACCTCACTCCTGTACTTACCTAAACAGATATAAATGGCTGGTTTTTAA GAAAAAAAA

SEO ID NO. 10

>11_TCC_25F2_M13F.TXT ,constant: -1, poly A: no ACCCTGGGAGAGAGTTTGAAGAAACCACAGCTGATGGCAGAAAAACTCAGACTGCTGCA ACTTTACAGATGGTGCATTGNGTCAGCATAGGAGTGAGATGGGGAAAGAAAGCACANTAA CAAGAAAATTGANAGATGNTAAATTAGTGNTGGAGTGTGTCATGAACAATGCACCTGT

SEO ID NO. 11

>25_TCC_50G5_M13F.TXT ,constant: 17, poly A: yes
TAGTGTGGAAGCATAGTGAACACACTGATTAGGTTATGGTTTAATGTTACAACAACTATT
TTTTAAGAAAAAACATGTTTTAGAAATTTGGTTTCAAGTGACATGTGTGAAAACAATATCG
ATACTACCATAGTGAGCCATGATTTTCTAAAAAAAAA

SEO ID NO. 12

>26_TCC_50G6_M13F.TXT ,constant: 17, poly A: yes
TAGTGTGGAAGCATAGTGAACACACTGATTAGGTTATGGTTTAATGTTACAACAACTATT
TTTTAAGAAAAACAAGTTTTAGAAATTTGGTTCAAGTGACATGTGTGAAAACAATATTGT
ATACTACCATAGTGAGCCATGATTTTCTAAAAAAAAA

SEO ID NO. 13

SEO ID NO. 14

SEO ID NO. 15

>38_TCC_56E11_M13F.TXT ,constant: -1, poly A: yes CTCTCCAGTTTGCACCTGTCCCCACCCTCCACTCAGCTGTCCTGCAGCAAACACTCCACC CTCCACCTTCCATTTTCCCCCACTACTGCAGCACCTCCAGGCCTGTTGCTATAGAGCCTA CCTGATGTCAATAAACAACAGCTGAAGCAAAAAAAAA

SEO ID NO. 16

SEO ID NO. 17

>53_TCC_79G2_M13F_E07_054.abl.TXT ,constant: 16, poly A: no TGTCCGTCTTCACCCATCCCCAAGCCTACTAGAGCAAGAAACCAGTTGTAATATAAAATG CACTGCCCTACTGTTGTTATGACTACCGTTACCTACTGTTGTCATTGTTATTACAGCTAT GGCCACTATTATTAAAGAGCTGTGTAACATCAAAAAAA

SEO ID NO. 18

SEO ID NO. 19

>35_TCC_21D6_M13F_C05_037.ab1.fa TIME: Wed Aug 9 12:48:31 2000 trimming information: raw_sequence:889 (high quality:34-340) sequence:95-456 [length:362] CTTTGACGTGGAGAGGACTCCTGCAATAACTTCATCTATGGAGGCTGCC GGGGCAATAAGAACAGCTACCGCTCTGAGGAGGCCTGCATGCTCCGCTGC TTCCGCCAGCAGGAGAATCCTCCCCTTGCCCCTTGGCTCAAAGGTGGTGCT TCTGGCGGGGCTGTTCGTGATGGTGTTGATCCTCTTCCTGGGAGCCTCCA TGGTCTACCTGATCCGGGTGGCACGGAGGAACCAGGAGCGTGCCCTGCGC ACCGTCTGGAGCTCCGAAGAGGAGCAGCTGGTGAAGAACACATA TGTCCTGTGACCGCCCTGTCGCCAAGAGGACCTGGNGAAAGGGAGGGAGA CTATGTGTGAGC

SEO ID NO. 20

>46_TCC_27H5_M13F_F06_058.abl.fa TIME: Wed Aug 9 12:48:35 2000 trimming information: raw_sequence:892 (high quality:169-406) sequence:170-287 [length:118] AAAAAGAGTAAAACACTTTCAGTTTCTCCCCTTTAGCCCCTAAAACAACA TCTTACAGTCTGGATCTACCTATACAGTCCTACATTAGCTTCTA AAATATTTGTCAGGAGGG